

From: Hoegh, Andrew [andrew.hoegh@montana.edu]
Sent: Thursday, March 11, 2021 1:16 PM
To: Manuel Ruiz [m.ruiz.aravena@gmail.com]
CC: Plowright, Raina [raina.plowright@montana.edu]; Clif McKee [clifton.mckee@gmail.com]; Alison Peel [a.peel@griffith.edu.au]
Subject: Re: [Viruses] Special Issue: Bats and Coronaviruses- Invitations

To finish this thought.... I'm not sure that would be a bats + coronavirus application. For now the manuscript says "Under Review," so maybe it will get sent out for peer review at MEE.

This particularly special issue has a July 31 deadline, so we can still "wait and see."

Andy

On Mar 11, 2021, at 1:13 PM, Andy Hoegh <andrew.hoegh@montana.edu> wrote:

Hi Manuel,

Yes, I think we could set up a scenario where other's data could fill in the narrative for this method.

On Mar 9, 2021, at 10:06 AM, Manuel Ruiz <m.ruiz.aravena@gmail.com> wrote:

Andrew... any chance of using the method with data from other papers?

I imagine we could have a look at papers with positive/negative samples and simulate pooling of samples, etc... (?)

Manuel Ruiz Aravena
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On Tue, Mar 9, 2021 at 10:02 AM Plowright, Raina

<raina.plowright@montana.edu> wrote:

Hi Andy, yes exactly. Don't want to spoil the big papers but this could be a good one for some scraps!

On Mar 9, 2021, at 9:18 AM, Hoegh, Andrew

<andrew.hoegh@montana.edu> wrote:

Hi Raina and Ali,

I'll defer to your perspective on this. I'm happy to summarize the sampling and/or analysis from the preliminary Australian CoV work. If the sampling-data integration gets rejected at MEE (which I would bet that it does), this could be reworked and use a small amount of the data (maybe even a single site - time). However, as you are well aware, the Australian CoV dataset is going to be a gold mine and we want to avoid spoiling a big paper from that effort.

Andy

On Mar 8, 2021, at 9:20 AM,

Plowright, Raina

<raina.plowright@montana.edu>

wrote:

Hi all,

I just received this. Peng Zhou is well known in the bat virus world (Wuhan Institute of Virology) and this could be a great opportunity to get some early Australian CoV work published if we have some small background papers we need to get out. Alternatively, Manuel and Clif, if we have some left-overs from the review.

Viruses has IF around 3 so its not a huge deal but papers in special issues also tend to get read and cited.

Raina

From: Allison Yang/MDPI
<allison.yang@mdpi.com>

Date: Monday, March 8,
2021 at 12:56 AM

To: Plowright, Raina
<raina.plowright@montana.edu>

Cc: 周
鹏 <peng.zhou@wh.iov.cn>
>, viruses@mdpi.com <viruses@mdpi.com>

Subject: [Viruses] Special
Issue: Bats and
Coronaviruses-
Invitations

Dear Dr. Plowright,

I would like to invite you to contribute a comprehensive review paper or full research article to the Special Issue "Bats and Coronaviruses" of *Viruses* (IF 3.816). *Viruses* is an open access journal indexed by Web of Science, PubMed, and other relevant databases. As Guest Editor, our aim for the Special Issue is to understand the relationship between bats and coronaviruses, and make efforts on the discovery and active surveillance of bat coronaviruses.

We would appreciate hearing from you soon regarding whether you could submit a contribution related to this field. Of course, you are welcome to invite any co-authors you consider appropriate. In case you are

interested, please let us know the specifics of the topic you would like to cover by providing a preliminary concept or tentative title or preliminary abstract in order to assist us with preparations for the Special Issue. A waiver of 30% of the publication charge would apply as a special offer for you.

We encourage authors to send a short abstract or tentative title to me and the Editorial Office (viruses@mdpi.com or allison.yang@mdpi.com) in advance. You may send your manuscript at any time prior to the deadline of **31 July 2021**.

More information may be found at the Special Issue website:
https://www.mdpi.com/journal/viruses/special_issues/coronaviruses_bats_2021

Please do not hesitate to contact us if you need any further information.

We look forward to hearing from you soon.

Best wishes,
Peng Zhou, PhD, Professor

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